Sequence: 12162674

Samples:	17910	Quality:	0 - 9
Bases:	1496		10 - 19
Average spacing:	12.0		20 - 29
Average quality >=	10: 111, 20: 97, 30: 1049		>= 30

GGGGGGAAAATTCCTAACACCGAGTTCGTGAAGGTGAAGGGCCTCCACTTCAGCCAGGAGGACGCTCCAGATGAAATGGGTAAGTA

CATCAAGAGCTTCGTGGAGCGCGTGCTGAAGAACGAGCAGTAATTCTAGGCGATCGCTCGAGTCAGCAGTGAACTCCTAAGTC

AAACCTATCAAATCCTTGTAATGAACCTGTAACCATTCGTCTTTTTATAATGAGATTTCTTAAATTTGGTAAGAGATTTAGC

ATTTGACTCAGCTCCCTGGTCAGACTTGTTTTTTGAGGACCTGGACAGCGTTCCATCCCTGGGAGTTAAAACCTGACATAGGA

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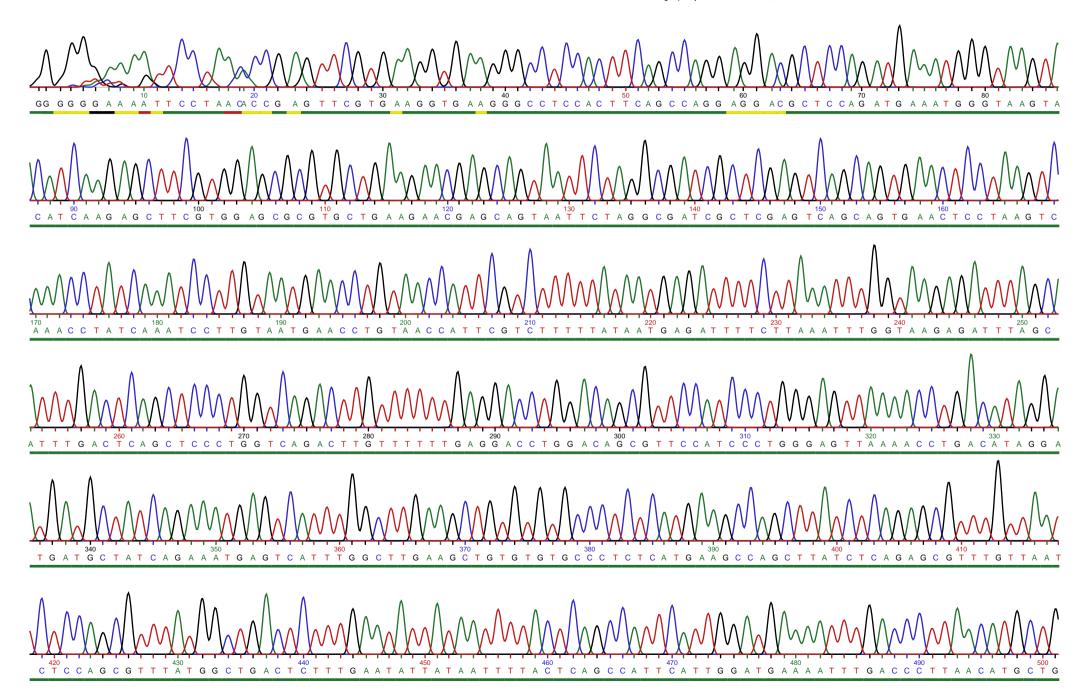CTCCAGCCGTTTATGGCTGACTCTTTGAATATTATAATTTTACTCAGCCATTCATTGGATGAAAATTTGACCCTTAACATGCTG

Sequence: 12162674

Samples: 17910
Bases: 1496
Average spacing: 12.0
Average quality >= 10: 111, 20: 97, 30: 1049

Quality: 0 - 9
10 - 19
20 - 29
>= 30

CTAAAGAAACAAATAGAAACTTAGTGCACGTCCATCTATTAAGTCAGGCACTGAGTCTGTGGCTCTCGTCTTCTGACCCGGCA

GTACAATTCTGTCTTGACATTTCTATACCAGTGCCAACACCACTCCTGGTCAAGACAGACAGATCCTTAACCAGTGCATAGCG

CATGCACTGGCTCCCTCTGCAGTCAGTTGGTCACCCCCCAACCCCCCTGTGCAACCCAGGTAGTCACTCCTCTACTGTCCCT

GGGAATTAATCTTGTCTGTTCTATTATTTCATTTAAGTTGAATGAGATGAGTTATACTTTTGTGTTGAAGTTTTATTTTTAGC

TTAACATGTTTTGACATGTGTGCTGTTGTATTTATCAATAGATCGTTTCTTTTTATTGGTGAATAGTATTCCTTATAGTAT

ACTTCTTTATACTAAGTAATATACTACTATTGGCCAATCTGTGTATCTGTTGATAGATACTCGGGATATTTCAGGGGGGGGGGAAAGG

Sequence: 12162674

Samples: 17910
Bases: 1496
Average spacing: 12.0
Average quality >= 10: 111, 20: 97, 30: 1049
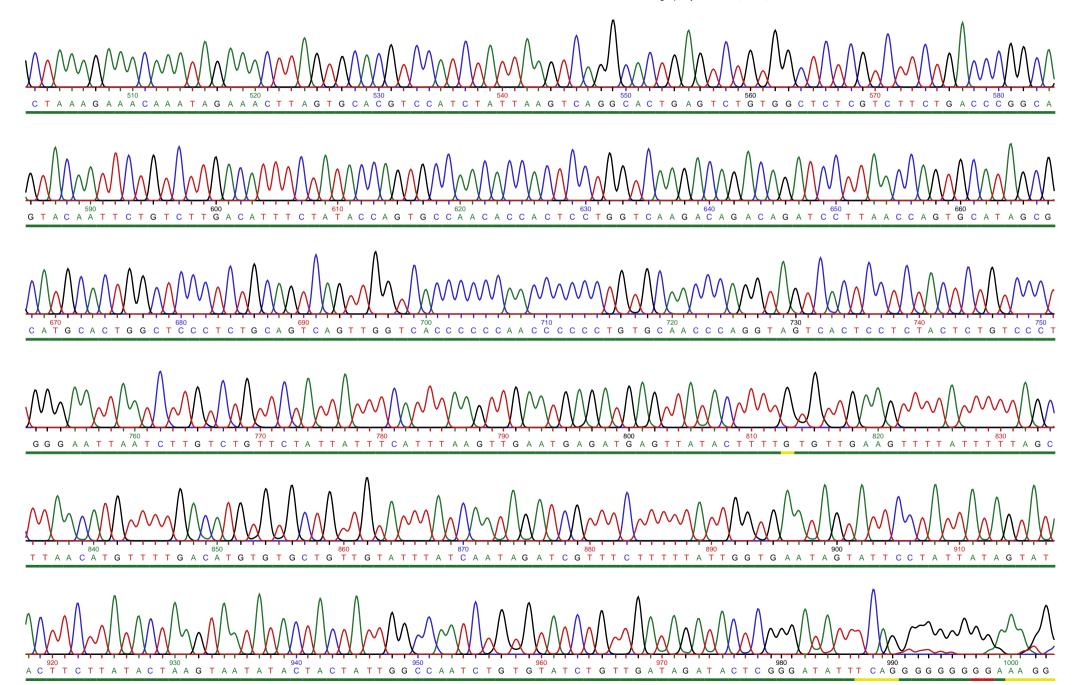
Quality: 0 - 9
10 - 19
20 - 29
>= 30

03.09.2019

1010 1020 1030 1040 1050 1060 1070 1080
C G G C C G C T G G C C G C A A T A A A A T A T C T T T A T T T C A T A C A T C C G G G T G GG T G G G T T T T T G G G T G A G G A T C T A A A T G A G T C T T C G G

1090 1100 1110 1120 1130 1140 1150 1160
A C C T C T C G G G G G C G C T T A A C G G G G G T A A A G G T T T G T C T G A C C G G G G G G G G G G G G G A G G G A A G A A C A C C C C C C T T T C C G G

1170 1180 1190 1200 1210 1220 1230 1240 1250
A G C G C G C G G G G T T G G G G G T T G G G G G G G G C A C C G G G A C A C C A A A A A A A C G C C A G A T C C C C T T A T A A C C C C C C C C C C C C C G G G

1260 1270 1280 1290 1300 1310 1320 1330
G G G G G G G G G T T T G T G C G G C C G G G G G G A A A A G G C G G G C C C C A A C C A C G C G G G G C G G C G G C C C C C A A G T G G T A A C T T T T T T T T T T

1340 1350 1360 1370 1380 1390 1400 1410
T T C C C A C A C C C C C G G G G A G C C C C C C C G C G C A G G A C T T T T T T T C T T C G C T C C C G C G G G G A A C A T T A A G A A G G G G C G T

1420 1430 1440 1450 1460 1470 1480 1490
G G A A C C C C C C A C C C C G C G G G A G A T A T T A A A A A A T A T T C C T G A A A A A A A A G T G A C T T T A T T T A T A G A A A A C A C T T G